## Experimental Results: Random digits and hypothesis testing

## 48 marks

In this question, we will explore the data collected in class.

You will need to download the data class\_data.csv from the assignment website and save it somewhere. Supposing the data to have been saved in a directory dataDirectory, read it into R as

Having loaded the data, you might have a look at its contents using any of the standard functions:

```
# just print it
data
# view its contents as a spreadsheet (in RStudio)
View(data)
# or, look at its data structure
str(data)
```

You will find that it is a data.frame with variables

```
names(data)
## [1] "random_digit" "student_digit" "green_card1" "green_card2"
```

```
## [1] "random_digit" "student_digit" "green_cardi" "green_card2" ## [5] "red_card1" "red_card2"
```

Each row contains one student's answer to the questions associated with these names.

Here we will only consider the results on the digits (0-9) obtained in class.

Recall how the data on the digits 0, 1, ..., 9 were collected.

Each person was first asked to write the words "random digit" on a card. They were then given a few seconds to think up a single *random* digit from 0 to 9 and then record it on the card.

On the other side of the card, each person then wrote "student digit" and below it recorded the *last* digit of their student id number.

These two digits provide the values for the variables random\_digit and student\_digit appearing in the data set data above.

1. Suppose a digit, d, is generated as a realization from a random variable D which is uniformly distributed on the digits  $\{0, 1, 2, 3, \dots, 8, 9\}$ . That is, for any value  $d \in \{0, 1, 2, 3, \dots, 8, 9\}$  we have

$$Pr(D=d) = \frac{1}{10}.$$

a. \*(1 mark)\*Determine the expectation \$E(D)\$.

$$E(D) = \sum_{D=0}^{9} D \cdot P(D) = \sum_{D=0}^{9} \frac{D}{10} = 4.5$$

b. \*(2 marks)\* Determine the expectation  $E(D^2)$  and hence the standard deviation D(D).

$$E(D) = \sum_{D=0}^{9} D^2 \cdot P(D) = \sum_{D=0}^{9} \frac{D^2}{10} = 28.5$$

$$Var(D) = E(D^2) - E(D)^2 = 28.5 - 4.5^2 = 8.25$$

$$SD(D) = \sqrt{(Var(D))} = \sqrt{8.25} = 2.872281$$

c. (1 mark) Determine the median of D.

$$median(D) = 4.5$$

d. Suppose we consider the random variable C which for some fixed value  $d \in A$  in 0, 1, 2, 3, 1

$$C = \begin{cases} 1 & \text{when } D = d \\ 0 & \text{when } D \neq d \end{cases}$$

which implies

$$Pr(C=1) = Pr(D=d) = \frac{1}{10}$$

and

$$Pr(C=0) = Pr(D \neq d) = 1 - Pr(D=d) = \frac{9}{10}$$

Suppose we have  $C_1$ ,  $C_2$ , dots, dots,

Let  $X = \sum_{i=1}^n C_i$ .

- i. \*(1 mark)\* What is the name of the probability distribution of \$X\$?
   binomial distribution
- ii. \*(1 mark)\* Hence, write down an expression Pr(X = x) for  $x \in \{0, 1, \ldots, n\}$ .

$$Pr(X=x) = \binom{n}{k} \left(\frac{1}{10}\right)^k \left(1 - \frac{1}{10}\right)^{n-k}$$

iii. \*(1 mark)\* Hence, write down an expression \$E(X)\$.

$$E\left(X\right) = np = \frac{n}{10}$$

e. Using the `data` from class, for \*\*each\*\* of the variables `random\_digit` and `student\_digit`, calcu

i. \*(2 marks)\* sample average,

```
s_digit <- data$student_digit
r_digit <- data$random_digit
s_digit_mean <- mean(s_digit)
r_digit_mean <- mean(r_digit)
print(s_digit_mean)</pre>
```

## [1] 4.595238

```
print(r_digit_mean)
```

- ## [1] 5.5
  - The mean of student\_digit is 4.595238
  - The mean of random\_digit is 5.5

ii. \*(2 marks)\* sample standard deviation,

```
s_digit_sd <- sd(s_digit)
r_digit_sd <- sd(r_digit)
print(s_digit_sd)</pre>
```

## [1] 3.052864

```
print(r_digit_sd)
```

- ## [1] 2.778401
  - The standard deviation of student\_digit is 3.052864
  - The standard deviation of random\_digit is 2.778401

iii. \*(2 marks)\* sample median,

```
s_digit_median <- median(s_digit)
r_digit_median <- median(r_digit)
print(s_digit_median)</pre>
```

## [1] 5

```
print(r_digit_median)
```

- ## [1] 7
  - The median of student\_digit is 5
  - The median of random\_digit is 7

iv. \*(3 marks)\* and compare these to the corresponding theoretical values from the distribution of

Varname	Theoretical	student_digit	random_digit
mean	4.5	4.595238	5.5
median	4.5	5	7
standard deviation	2.872281	3.052864	2.778401

- \* student\_digit sample has larger mean, median, and standard deviation than theoretical value
- \* random\_digit sample has larger mean, and median than theoretical results.
- \* But its standard deviation is smaller than theoretical value
- v. \*(1 mark)\* Which of `random\_digit` and `student\_digit` have sample values closer to the theoreti
  - student\_digit have sample values closer to the theoretical
     values because the median and mean of student\_digit
     is closes to theroretical values. student\_digit and random\_digit
     have standards deviation are all close to the theoretical sd.
- -- Student\_digit is more close to theoretical(uniform distribution)
- f. \*(3 marks)\* Calculate Pr(X = x) when n = 42 and x = 0, 5, and 10.

$$\begin{split} Pr(X=0) &= \binom{42}{0} \left(\frac{1}{10}\right)^0 \left(\frac{9}{10}\right)^{42-0} = 0.01197251518 \\ Pr(X=5) &= \binom{42}{5} \left(\frac{1}{10}\right)^5 \left(\frac{9}{10}\right)^{42-5} = 0.17247769726 \\ Pr(X=10) &= \binom{42}{10} \left(\frac{1}{10}\right)^{10} \left(\frac{9}{10}\right)^{42-10} = 0.00505246993 \end{split}$$

2. We are interested testing the hypothesis

H: the observed digits  $d_1$ ,  $d_2$ , ...,  $d_n$  are independent realizations of a random variable D uniformly distributed on the digits  $\{0, 1, 2, ..., 9\}$ .

In particular, we are interested in testing this hypothesis for each of two samples student\_digit and random\_digit.

- a. (4 marks) The function stem() is a simple way to get a quick picture (a "stem and leaf plot") of the distribution of a set of digits. Use stem() to construct a picture of each of the following:
  - i. the values of student\_digit,

##

3 | 0000

```
stem(s_digit)
##
##
     The decimal point is at the |
##
##
     0 | 000000
##
     1 | 0000
##
     2 | 000
##
     3 | 0
     4 | 000000
##
##
     5 | 000
##
     6 | 0000000
##
     7 | 0
##
     8 | 0000000
     9 | 0000
##
    ii. the values of `random_digit`,
stem(r_digit)
##
##
     The decimal point is at the |
##
##
     0 | 000
     1 | 0
##
##
     2 | 000
##
     3 | 000000
##
     4 | 00
     5 | 00
##
     6 | 000
##
     7 | 00000000000
##
     8 | 0000
##
     9 | 000000
##
    iii. and, for comparison, a sample of the same size from a uniform distribution on the digits using
        the function `sample()`.
uniform_sample <- sample(1:10, replace = TRUE,40)</pre>
stem(uniform_sample)
##
##
     The decimal point is at the |
##
##
      1 | 0000
      2 | 00
##
```

```
## 4 | 000000

## 5 | 000

## 6 | 00000

## 7 | 00

## 8 | 000

## 9 | 0000000

## 10 | 0000
```

Which of `student\_digit` or `random\_digit` looks more like it might have come from a Uniform on the

- student\_digit looks more like it might have come from a Uniform on the digits becaus the leng

b. A more formal way to assess whether a sample of values appear to come from a hypothesized distributi

$$\chi^2 = \sum_{i=1}^{m} \frac{(o_i - e_i)^2}{e_i}$$

where \$o\_i\$ is the observed number of values in the \$i\$th "cell", \$e\_i\$ is the expected number to form our case, the cells are the 10 different possible digits (so \$m = 10\$) and \$o\_i\$ is the number of \$\Chi^2\$ is a discrepancy measure which is larger whenever the \$o\_i\$ are relatively far from their form the hypothesized model is true, then the distribution of \$\Chi^2\$ can usually be approximated as

i. \*(4 marks)\* Write a function `count\_digits()` which takes a vector of digits `d` and returns a n

```
count_digits <- function (d) {</pre>
  len<- length(d)</pre>
  digits \leftarrow seq(0,len-1,1)
  return(sapply(digits,function(x) sum(d==x))[1:10])
my_digits \leftarrow c(0, 1, 3, 4, 7, 1, 4, 9, 7, 4)
print("my_function returns:")
## [1] "my_function returns:"
count_digits(my_digits)
## [1] 1 2 0 1 3 0 0 2 0 1
# would return a vector equal to
print("it should be: ")
## [1] "it should be: "
c(1, 2, 0, 1, 3, 0, 0, 2, 0, 1)
## [1] 1 2 0 1 3 0 0 2 0 1
        # for example if
        my_digits \leftarrow c(0, 1, 3, 4, 7, 1, 4, 9, 7, 4)
        # then
        count_digits(my_digits)
```

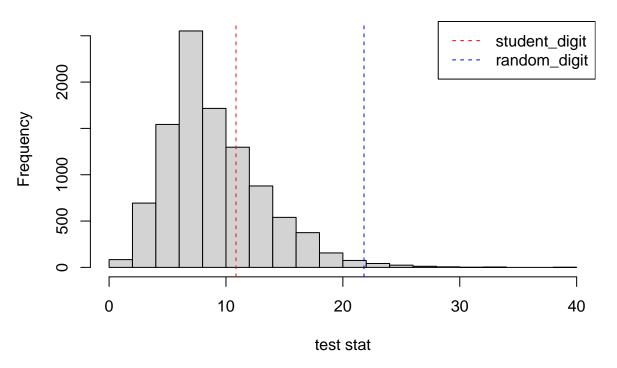
```
# would return a vector equal to
        c(1, 2, 0, 1, 3, 0, 0, 2, 0, 1)
       Note, that we will assume that `d` will be an integer vector containing
       only values in \{0, 1, 2, \ldots, 9\}.
       No error checking is required for now.
    ii. *(2 marks)* Demonstrate your function on the digits of the variable `student_digit` and on the
count_digits(s_digit)
## [1] 6 4 3 1 6 3 7 1 7 4
count_digits(r_digit)
## [1] 3 1 3 6 2 2 3 12 4 6
    iii. *(4 marks)* Write the function `Pearson_chi_sq(observed, expected)` which calculates $\Chi^2$
            Again, for expediency it will be assumed that both `observed` and `expected` vectors contain
            However, you should check that the lengths of `observed` and `expected` match and stop if to
Pearson_chi_sq <- function (observed,
                        expected = sum(observed)/length(observed)) {
  if(length(expected) == 1){
    expected = rep(expected,length((observed)))
  if(length(observed) != length(expected)){
   print("Lengths do not match")
    return()
  values <- mapply(function(x,y)\{(x-y)^2/y\}, observed, expected)
  return(sum(values))
Pearson_chi_sq(count_digits(data$student_digit))
## [1] 10.85714
Pearson_chi_sq(count_digits(data$random_digit))
## [1] 21.80952
        - The Pearson_chi_sq
    iv. *(2 marks)* Check your function by comparing the values of `Pearson_chi_sq(observed)` to resul
        `count_digits(data$student_digit)` and `count_digits(data$random_digit)` in turn.
chisq.test(count_digits(data$student_digit))$statistic
## Warning in chisq.test(count_digits(data$student_digit)): Chi-squared
## approximation may be incorrect
```

```
## 10.85714
chisq.test(count_digits(data$random_digit))$statistic
## Warning in chisq.test(count_digits(data$random_digit)): Chi-squared
## approximation may be incorrect
## X-squared
## 21.80952
    v. *(2 marks)* Using the function `pchisq()` calculate the $p$-value testing the uniformity hypothe
    Show that your -values agree with those produced by chisq.test(observed)$p.value for the counts of
pchisq(Pearson_chi_sq(count_digits(data$student_digit)),9,lower.tail = FALSE)
## [1] 0.2856284
pchisq(Pearson_chi_sq(count_digits(data$random_digit)),9,lower.tail = FALSE)
## [1] 0.009502653
chisq.test(count_digits(data$student_digit))$p.value
## Warning in chisq.test(count_digits(data$student_digit)): Chi-squared
## approximation may be incorrect
## [1] 0.2856284
chisq.test(count_digits(data$random_digit))$p.value
## Warning in chisq.test(count_digits(data$random_digit)): Chi-squared
## approximation may be incorrect
## [1] 0.009502653
    vi. *(3 marks)* Rather than depend upon the validity of the $\chi^2_k$ approximation, we could *sim
        Using the function `sapply()` and the function `sample()`, together with your functions `Pearson
        That is, write
get_chisqs <- function (n, B = 1000) {</pre>
  count_list <- list()</pre>
  for (i in 1:B){
    my_sample <- sample(0:9, n, replace = TRUE)</pre>
    count_list[[i]] <- count_digits(my_sample)</pre>
  chisq_list<-sapply(1:B, function(x)</pre>
  Pearson_chi_sq(count_list[[x]]))
  return(chisq_list)
}
n <- nrow(data)
results <- get_chisqs(n = n, B = 1000)
#print(results)
```

## X-squared

vii. \*(3 marks)\* Use your function `get\_chisqs()` to get `B = 10000` independent pseudo-random rea
That is, execute the following (N.B. in RMarkdown change header to eval = TRUE):

## Simulated Pearson test null distribution



To this histogram, add a vertical line in "red" where the corresponding statistics you calc

Put a legend in the top right that identifies the lines.

Based on this histogram, which collection of digits seem less likely to have been generated

- Random digit is lesslikely to have been generated as a random sample.
- The larger is X2, the greater is the evidence against the model.
- random\_digit has large X2 value.

viii. \*(2 marks)\* The simulated distributions can be used to calculate approximate \$p\$-values by si

Calculate these two \$p\$-values using the simulated test null distribution given by the vect

```
chi_sq_student <- Pearson_chi_sq(count_digits(data$student_digit))
p_value_student <- sum(chisq_stats >= chi_sq_student)/length(chisq_stats)
print(p_value_student)

## [1] 0.3013

chi_sq_random <- Pearson_chi_sq(count_digits(data$random_digit))
p_value_random <- sum(chisq_stats >= chi_sq_random)/length(chisq_stats)
print(p_value_random)

## [1] 0.0107
```

- ix. \*(2 marks)\* What do you conclude about the two hypotheses?
  - For student digit, the p-value > 0.05, which means that the null hypothesis can be accepted.
  - For random digit, the p-value < 0.05, which means that the null1 hypothesis should be rejected.
  - student\_digit is randomly selected, the random\_digit is NOT randomly selected.